

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/033, 129B
Source: FW16
Date Processed by STIC: 1-3-05

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 01/03/2005
 PATENT APPLICATION: US/10/033,129B TIME: 14:56:25

Input Set : A:\8325-2001-30seqlistcomplete.txt
 Output Set: N:\CRF4\01032005\J033129B.raw

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3 <110> APPLICANT: CHOO, Yen
4      KLUG, Aaron
5      SANCHEZ-GARCIA, Isidro
7 <120> TITLE OF INVENTION: BINDING PROTEINS FOR RECOGNITION OF DNA
9 <130> FILE REFERENCE: 8325-2001.30 (G1-US3)
11 <140> CURRENT APPLICATION NUMBER: 10/033,129B
12 <141> CURRENT FILING DATE: 2001-12-27
14 <150> PRIOR APPLICATION NUMBER: 08/793,408
15 <151> PRIOR FILING DATE: 1997-06-02
17 <150> PRIOR APPLICATION NUMBER: PCT/GB95/01949
18 <151> PRIOR FILING DATE: 1995-08-17
20 <160> NUMBER OF SEQ ID NOS: 125
22 <170> SOFTWARE: PatentIn version 3.2
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 60
26 <212> TYPE: DNA
27 <213> ORGANISM: Artificial
29 <220> FEATURE:
30 <223> OTHER INFORMATION: oligonucleotide
32 <400> SEQUENCE: 1
33 ctcctgcagt tggacctgtg ccatggccgg ctgggcccga tagaatggaa caactaaagc      60
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 92
38 <212> TYPE: PRT
39 <213> ORGANISM: Artificial
41 <220> FEATURE:
42 <223> OTHER INFORMATION: zinc finger protein used in phage display library
45 <220> FEATURE:
46 <221> NAME/KEY: MISC_FEATURE
47 <222> LOCATION: (48)..(51)
48 <223> OTHER INFORMATION: Xaa = any amino acid
50 <220> FEATURE:
51 <221> NAME/KEY: MISC_FEATURE
52 <222> LOCATION: (53)..(54)
53 <223> OTHER INFORMATION: Xaa = any amino acid
55 <220> FEATURE:
56 <221> NAME/KEY: MISC_FEATURE
57 <222> LOCATION: (56)..(56)
58 <223> OTHER INFORMATION: Xaa = any amino acid
60 <220> FEATURE:
61 <221> NAME/KEY: MISC_FEATURE
62 <222> LOCATION: (57)..(57)
63 <223> OTHER INFORMATION: Xaa = Arg or Lys

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65 <400> SEQUENCE: 2
67 Met Ala Glu Glu Arg Pro Tyr Ala Cys Pro Val Glu Ser Cys Asp Arg
68 1 5 10 15
71 Arg Phe Ser Arg Ser Asp Glu Leu Thr Arg His Thr Arg Ile His Thr
72 20 25 30
W--> 75 Gly Gln Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Xaa
76 35 40 45
79 Xaa Xaa Xaa Leu Xaa Xaa His Xaa Xaa Thr His Thr Gly Glu Lys Pro
80 50 55 60
83 Phe Ala Cys Asp Ile Cys Gly Arg Lys Phe Ala Arg Ser Asp Glu Arg
84 65 70 75 80
87 Lys Arg His Thr Lys Ile His Leu Arg Gln Lys Asp
88 85 90
91 <210> SEQ ID NO: 3
92 <211> LENGTH: 26
93 <212> TYPE: DNA
94 <213> ORGANISM: Artificial
96 <220> FEATURE:
97 <223> OTHER INFORMATION: oligonucleotide
99 <400> SEQUENCE: 3
100 tatgacttgg atgggagacc gcctgg 26
103 <210> SEQ ID NO: 4
104 <211> LENGTH: 28
105 <212> TYPE: DNA
106 <213> ORGANISM: Artificial
108 <220> FEATURE:
109 <223> OTHER INFORMATION: oligonucleotide
111 <400> SEQUENCE: 4
112 aattccaggc ggtctcccat ccaagtca 28
115 <210> SEQ ID NO: 5
116 <211> LENGTH: 21
117 <212> TYPE: DNA
118 <213> ORGANISM: Artificial
120 <220> FEATURE:
121 <223> OTHER INFORMATION: oligonucleotide
123 <400> SEQUENCE: 5
124 tatatagcgt gggcgatatat a 21
127 <210> SEQ ID NO: 6
128 <211> LENGTH: 24
129 <212> TYPE: DNA
130 <213> ORGANISM: Artificial
132 <220> FEATURE:
133 <223> OTHER INFORMATION: oligonucleotide
135 <400> SEQUENCE: 6
136 gcgtatatac gcccacgcta tata 24
139 <210> SEQ ID NO: 7
140 <211> LENGTH: 21
141 <212> TYPE: DNA
142 <213> ORGANISM: Artificial

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144 <220> FEATURE:
145 <223> OTHER INFORMATION: oligonucleotide
148 <220> FEATURE:
149 <221> NAME/KEY: misc_feature
150 <222> LOCATION: (10)..(12)
151 <223> OTHER INFORMATION: n is a, c, g, or t
153 <400> SEQUENCE: 7
W--> 154 tatatagcgn nngcgatat a 21
157 <210> SEQ ID NO: 8
158 <211> LENGTH: 24
159 <212> TYPE: DNA
160 <213> ORGANISM: Artificial
162 <220> FEATURE:
163 <223> OTHER INFORMATION: oligonucleotide
166 <220> FEATURE:
167 <221> NAME/KEY: misc_feature
168 <222> LOCATION: (13)..(15)
169 <223> OTHER INFORMATION: n is a, c, g, or t
171 <400> SEQUENCE: 8
W--> 172 gcgtatatagc gcnngcgcta tata 24
175 <210> SEQ ID NO: 9
176 <211> LENGTH: 33
177 <212> TYPE: DNA
178 <213> ORGANISM: Artificial
180 <220> FEATURE:
181 <223> OTHER INFORMATION: BCR-ABL
183 <400> SEQUENCE: 9
184 ttccatggag acgcagaagc ccttcagcgg cca 33
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 33
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial
192 <220> FEATURE:
193 <223> OTHER INFORMATION: BCR
195 <400> SEQUENCE: 10
196 ttccatggag acgcaggtga gttcctcacg cca 33
199 <210> SEQ ID NO: 11
200 <211> LENGTH: 33
201 <212> TYPE: DNA
202 <213> ORGANISM: Artificial
204 <220> FEATURE:
205 <223> OTHER INFORMATION: ABL
207 <400> SEQUENCE: 11
208 cccctttctc ttccagaagc ccttcagcgg cca 33
211 <210> SEQ ID NO: 12
212 <211> LENGTH: 33
213 <212> TYPE: PRT
214 <213> ORGANISM: Artificial
216 <220> FEATURE:

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217 <223> OTHER INFORMATION: 1A
 219 <400> SEQUENCE: 12
 221 Met Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe
 222 1 5 10 15
 225 Ser Asp Arg Ser Ser Leu Thr Arg His Thr Arg His Thr Gly Glu Lys
 226 20 25 30
 229 Pro
 233 <210> SEQ ID NO: 13
 234 <211> LENGTH: 33
 235 <212> TYPE: PRT
 236 <213> ORGANISM: Artificial
 238 <220> FEATURE:
 239 <223> OTHER INFORMATION: 1B
 241 <400> SEQUENCE: 13
 243 Met Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe
 244 1 5 10 15
 247 Ser Glu Arg Gly Thr Leu Ala Arg His Glu Lys His Thr Gly Glu Lys
 248 20 25 30
 251 Pro
 255 <210> SEQ ID NO: 14
 256 <211> LENGTH: 27
 257 <212> TYPE: PRT
 258 <213> ORGANISM: Artificial
 260 <220> FEATURE:
 261 <223> OTHER INFORMATION: 2A
 263 <400> SEQUENCE: 14
 265 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Gln Gly Gly Asn Leu
 266 1 5 10 15
 269 Val Arg His Leu Arg His Thr Gly Glu Lys Pro
 270 20 25
 273 <210> SEQ ID NO: 15
 274 <211> LENGTH: 26
 275 <212> TYPE: PRT
 276 <213> ORGANISM: Artificial
 278 <220> FEATURE:
 279 <223> OTHER INFORMATION: 3A
 281 <400> SEQUENCE: 15
 283 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Gln Ala Gln Thr Leu
 284 1 5 10 15
 287 Gln Arg His Leu Lys His Thr Gly Glu Lys
 288 20 25
 291 <210> SEQ ID NO: 16
 292 <211> LENGTH: 26
 293 <212> TYPE: PRT
 294 <213> ORGANISM: Artificial
 296 <220> FEATURE:
 297 <223> OTHER INFORMATION: 3B
 299 <400> SEQUENCE: 16
 301 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Gln Ala Ala Thr Leu

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Input Set : A:\8325-2001-30seqlistcomplete.txt

Output Set: N:\CRF4\01032005\J033129B.raw

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302 1          5          10          15
305 Gln Arg His Leu Lys His Thr Gly Glu Lys
306          20          25
309 <210> SEQ ID NO: 17
310 <211> LENGTH: 26
311 <212> TYPE: PRT
312 <213> ORGANISM: Artificial
314 <220> FEATURE:
315 <223> OTHER INFORMATION: 3C
317 <400> SEQUENCE: 17
319 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Gln Ala Gln Asp Leu
320 1          5          10          15
323 Gln Arg His Leu Lys His Thr Gly Glu Lys
324          20          25
327 <210> SEQ ID NO: 18
328 <211> LENGTH: 89
329 <212> TYPE: PRT
330 <213> ORGANISM: Artificial
332 <220> FEATURE:
333 <223> OTHER INFORMATION: zinc finger polypeptide
335 <400> SEQUENCE: 18
337 Met Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe
338 1          5          10          15
341 Ser Asp Arg Ser Ser Leu Thr Arg His Thr Arg Thr His Thr Gly Glu
342          20          25          30
345 Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Asp Arg Ser
346          35          40          45
349 His Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys Pro Phe Gln
350          50          55          60
353 Cys Arg Ile Cys Met Arg Asn Phe Ser Asp Arg Ser Asn Leu Thr Arg
354 65          70          75          80
357 His Thr Arg Thr His Thr Gly Glu Lys
358          85
361 <210> SEQ ID NO: 19
362 <211> LENGTH: 10
363 <212> TYPE: PRT
364 <213> ORGANISM: Artificial
366 <220> FEATURE:
367 <223> OTHER INFORMATION: zinc finger binding motif
369 <400> SEQUENCE: 19
371 Arg Ser Asp His Leu Thr Thr His Ile Arg
372 1          5          10
374 <210> SEQ ID NO: 20
375 <211> LENGTH: 10
376 <212> TYPE: PRT
377 <213> ORGANISM: Artificial
379 <220> FEATURE:
380 <223> OTHER INFORMATION: zinc finger binding motif
382 <400> SEQUENCE: 20

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/03/2005
PATENT APPLICATION: US/10/033,129B TIME: 14:56:26

Input Set : A:\8325-2001-30seqlistcomplete.txt
Output Set: N:\CRF4\01032005\J033129B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 48,49,50,51,53,54,56,57
Seq#:7; N Pos. 10,11,12
Seq#:8; N Pos. 13,14,15

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27
Seq#:28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51
Seq#:52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75
Seq#:76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99
Seq#:100,101,102,103,104,105,106,107,108,109,110,111,112,113,114,115,116,117
Seq#:118,119,120,121,122,123,124,125

VERIFICATION SUMMARY

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Input Set : A:\8325-2001-30seqlistcomplete.txt

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L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:32

M:341 Repeated in SeqNo=2

L:154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0

L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0